



SEQUENCE LISTING

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<110> Cantor, Charles R.
Przetakiewicz, Marek
Smith, Cassandra L.
Sano, Takeshi

<120> Positional Sequencing by Hybridization

<130> 25491-2401G

<140> US 09/030,571
<141> 1998-02-24

<150> US 07/792,012
<151> 1992-11-06

<150> US 08/110,691
<151> 1993-08-23

<150> US 08/470,832
<151> 1995-06-06

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<220>
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<221> misc_feature
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<223> r is a or g
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gtcgacagt gacgctacca ynnnnrtggc ctagagctag c 41

<210> 16
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<210> 23
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<400> 26
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<210> 27
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<221> misc_feature
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<221> misc_feature
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<400> 27
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<400> 35
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<400> 36
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<211> 13
<212> DNA
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<223> Bbv I Recognition Sequence

<221> misc_feature
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<211> 17
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<223> Bbv I Recognition Sequence - Complment

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<223> n is a, c, t or g

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<223> Bgl I Recognition Sequence

<221> misc_feature
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<223> n is a, c, t or g

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<223> Bgl I Recognition Sequence - Complement

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<223> n is a, c, t or g

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gccnnnnnngg c 11

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<221> misc_feature
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<223> n is a, c, t or g

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ccannnnnnnt gg 12

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<221> misc_feature
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<211> 9
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<221> misc_feature
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cacnnnngtg 9

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<220>
<223> Dra III Recognition Sequence - Complement

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<210> 45
<211> 14
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<223> Fok I Recognition Sequence

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<210> 47
<211> 10
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<213> Artificial Sequence

<220>
<223> Hga I Recognition Sequence

<221> misc_feature
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gacgcnnnnn 10

<210> 48
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<223> Hga I Recognition Sequence - Complement

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ccannnnntg g 11

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<211> 11
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<221> misc_feature
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<223> n is a, c, t or g

<400> 50
ccannnnnntg g 11

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<221> misc_feature
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<220>
<223> SfaN I Recognition Sequence - Complement

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<223> n is a, c, t or g

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nnnnnnnnng atgc 14

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<400> 53
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<210> 54
<211> 13
<212> DNA
<213> Artificial Sequence

<220>
<223> Sif I Recognition Sequence - Complement

<221> misc_feature
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<223> n is a, c, t or g

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ggccnnnnng gcc 13

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